



52288B-SEQ-listing.txt  
SEQUENCE LISTING

<110> Hillyard, Jeanna  
Roberts, James  
Ye, Minwei

<120> Cotton Event PV-GHBK04 (757) and Compositions and Methods for Detection Thereof

<130> 38-21 (52288)B

<140> 09/990,659

<141> 2001-11-16

<150> US60/249,757

<151> 2000-11-17

<160> 21

<170> PatentIn version 3.0

<210> 1

<211> 20

<212> DNA

<213> artificial sequence

<220>

<221> misc\_feature

<222> (1)..(20)

<223> a 5' genome-insert junction nucleotide sequence which is/is complementary to a sequence diagnostic for nucleic acids derived from the cotton event 757 recombinant genome

<400> 1

gtttgcttgg acactgatag

20

<210> 2

<211> 20

<212> DNA

<213> artificial sequence

<220>

<221> misc\_feature

<222> (1)..(20)

<223> a 3' genome-insert junction nucleotide sequence which is/is complementary to a sequence diagnostic for nucleic acids derived from the cotton event 757 recombinant genome

<400> 2

aaaccctttc tggaaaaata

20

<210> 3

<211> 20

<212> DNA

<213> Gossypium hirsutum

<220>

<221> misc\_feature

<222> (1)..(20)

RECEIVED  
APR 08 2002  
TECH CENTER 1600/2900

## 52288B-SEQ-listing.txt

```

<400> 3
tggtctgtgg aaaaggaagg 20

<210> 4
<211> 20
<212> DNA
<213> Gossypium hirsutum

<220>
<221> misc_feature
<222> (1)..(20)

<400> 4
atgcctgcag gtcaattcaa 20

<210> 5
<211> 138
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(138)
<223> part of a 5' non-functional sequence inserted into the cotton genome in cotton event 757

<400> 5
acactgatag tttaaactga aggcgggaaa cgacaatctg atcccagctt gcatgcctgc 60
aggtcaattc aatattgtgg caggacattg ctacatgata cctcttagaa ttgttttagac 120
ttcagatcga tcttgtca 138

<210> 6
<211> 767
<212> DNA
<213> Gossypium hirsutum

<220>
<221> Unsure
<222> (1)..(767)
<223> 5' cotton (Gossypium hirsutum) genome sequence

<400> 6
gtcccggggg cttatcctgt attcatttgc acccacataa acagccaaat taaccaaacc 60
catattcaac tgaaactccc aaagccattc ctacttttagc ttttcacca ctaactcaaa 120
agaaaacact cacctagctt ctttgctttt tcttttggat tgttttagat ctacaaaaag 180
atgattcaag aactccttgg aggttcttct tgcttaaact ttggagggga gaggaagatc 240
tccatcaatg gaagcatttt ggaaggaacc cccacttctt ctccatcacc atcatcttct 300
tcttcttcgg cgacgacttc atcgaccact aattcatcga atccggagaa tcatcaccag 360
aatgtgaggt gcccaggtg tgattcctcc aacacaaagt tctgctatta caacaactac 420
aacctcactc agcctcgtca cttttgcaag acttgccgtc ggtattggac caaaggagga 480

```

52288B-SEQ-listing.txt

```

gctctcagaa acgttcctat tgggtggtggg tgtaggaaaa acaaaagcac tactggtggt      540
tcaacatctc tggggaaatc aacttcttcc aagatgaaaa cagtagtttc tgaaattgga      600
agatctgggt tcgatcatga gcttcagtct actccaattc ttgggacttc agcggccccag      660
acttcccata ttctatccaa tctaacctca atgagagcta ccctaaaccc taaccctaac      720
acattgtcta accctgttag tattaaggaa gaagtgaagt tgcttggt                      767

```

```

<210> 7
<211> 206
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(206)
<223> part of arbitrarily assigned 3' end DNA sequence inserted into th
      e cotton event 757 genome.

```

```

<400> 7
tgagggatca agccacagca gccactcga ctttctagcc gaccagacg agccaaggga      60
tcttttttga atgctgctcc gtcgtcaggc tttccgacgt ttgggtggtt gaacagaagt      120
cattatcgca cggaatgcc agcactcccg aggggaaccc tgtggttggc atgcacatac      180
aaatggacga acggataaac cctttc                                           206

```

```

<210> 8
<211> 307
<212> DNA
<213> Gossypium hirsutum

<220>
<221> Unsure
<222> (1)..(307)
<223> 3' cotton (Gossypium hirsutum) genome sequence

```

```

<400> 8
tggaaaaata atcaacacca cgtcaacaa caacagaata ataatggggt cttgttaggt      60
gaagttcaaa acacaggtat tcaagaactg tatcaaaggc tcaaatac atcaagttat      120
tactctgata cttcagcagt aattctaagc aatgtcgctt cttcttcac aacatccatt      180
ttggagtcag ctccagttgc tgggggagaa ttgggttact ggaatccggc attttcatca      240
tcgtggtctg atcttccaac aactaatggt gcatatcctt aaaataaccc tttaccttc      300
gtttaat                                                                307

```

```

<210> 9
<211> 26
<212> DNA
<213> artificial sequence

```

<220>  
 <221> misc\_feature  
 <222> (1)..(26)  
 <223> 5' cotton (Gossypium hirsutum) genome PCR primer

<400> 9  
 gagagagata ggcactaaag taagca

26

<210> 10  
 <211> 28  
 <212> DNA  
 <213> artificial sequence

<220>  
 <221> misc\_feature  
 <222> (1)..(28)  
 <223> 5' insert PCR primer

<400> 10  
 ttagacaaat tgtcacgggtc taccagaa

28

<210> 11  
 <211> 24  
 <212> DNA  
 <213> artificial sequence

<220>  
 <221> misc\_feature  
 <222> (1)..(24)  
 <223> 3' insert PCR primer

<400> 11  
 ttcccaacga tcaaggcgag ttac

24

<210> 12  
 <211> 27  
 <212> DNA  
 <213> artificial sequence

<220>  
 <221> misc\_feature  
 <222> (1)..(27)  
 <223> 3' cotton (Gossypium hirsutum) genome PCR primer

<400> 12  
 ttgatgcact tacgaaagaa gaaccga

27

<210> 13  
 <211> 905  
 <212> DNA  
 <213> artificial sequence

<220>  
 <221> misc\_feature  
 <222> (1)..(905)  
 <223> 5' cotton (Gossypium hirsutum) genome + insert sequence

## 52288B-SEQ-listing.txt

```

<400> 13
gtcccggggg cttatcctgt attcatttgc acccacataa acagccaaat taaccaaacc      60
catattcaac tgaaactccc aaagccattc ctacttttagc ttttcaccca ctaactcaaa      120
agaaaacact cacctagctt ctttgctttt tcttttggat tgttttagat ctacaaaaag      180
atgattcaag aactccttgg aggttcttct tgcttaaact ttggagggga gaggaagatc      240
tccatcaatg gaagcatttt ggaaggaacc cccacttctt ctccatcacc atcatcttct      300
tcttcttcgg cgacgacttc atcgaccact aattcatcga atccggagaa tcatcaccag      360
aatttgaggt gccccaggtg tgattcctcc aacacaaagt tctgctatta caacaactac      420
aacctcactc agcctcgtca cttttgcaag acttgccgtc ggtattggac caaaggagga      480
gctctcagaa acgttcctat tgggtggtggg tgtaggaaaa acaaaagcac tactggtggt      540
tcaacatctc tggggaaatc aacttcttcc aagatgaaaa cagtagtttc tgaaattgga      600
agatctgggt tcgatcatga gcttcagtct actccaattc tttggacttc agcggcccag      660
acttcccata ttctatccaa tctaacctca atgagagcta ccctaaaccc taaccctaac      720
acattgtcta accctgttag tattaaggaa gaagtgagtt tgcttggaca ctgatagttt      780
aaactgaagg cgggaaacga caatctgata ccagcttgca tgccctgcagg tcaattcaat      840
attgtggcag gacattgcta catgatacct cttagaattg tttagacttc agatcgatct      900
tgtca                                           905

```

```

<210> 14
<211> 513
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(513)
<223> 3' cotton (Gossypium hirsutum) genome + insert sequence

```

```

<400> 14
tgagggatca agccacagca gccactcga ccttctagcc gaccagacg agccaagggga      60
tctttttgga atgctgctcc gtcgtcaggc tttccgacgt ttgggtggtt gaacagaagt      120
cattatcgca cggaatgcc agcactcccg aggggaaccc tgtggttggc atgcacatac      180
aatggacga acggataaac cctttctgga aaaataatca acaccacgct caacaacaac      240
agaataataa tgggttcctt gtaggtgaag ttcaaaacac aggtattcaa gaactgtatc      300
aaaggctcaa atcatcatca agttattact ctgatacttc agcagtaatt ctaagcaatg      360
tcgcttcttc ttcatacaac tccattttgg agtcagctcc agttgctggg ggagaattgg      420
gttactggaa tccggcattt tcatcatcgt ggtctgatct tccaacaact aatggtgcat      480

```

atccttaaaa taacccttta cctttcgttt aat

513

<210> 15  
 <211> 4973  
 <212> DNA  
 <213> artificial sequence

<220>  
 <221> misc\_feature  
 <222> (1)..(4973)  
 <223> sequence of 5' flank to full-length cry1Ac coding region

<400> 15  
 cggcccagac ttcccatctt ctatccaatc taacctcaat gagagctacc ctaaacccta 60  
 accctaacac attgtctaac cctgttagta ttaaggaaga agtgagtttg cttggacact 120  
 gatagtttaa actgaaggcg ggaaacgaca atctgatccc agcttgcatg cctgcaggtc 180  
 aattcaatat tgtggcagga cattgctaca tgatacctct tagaattggt tagacttcag 240  
 atcgatcttg tcagtctgaa agacccaaaa acaaatgcaa tttcttttct ggtagaccgt 300  
 gacaatttgt ctaagatgta tctgatttaa tgccttttgt atataatata ctcatctaata 360  
 ctagttaatt tagcttcaga gtaaattact tcagcatatt tatacgtgcc aagtgccaac 420  
 catatcaaata tagctaagca gacagttgaa gtacacaaaa caaaagcatc atatgctgat 480  
 ttattttattc atagatggag ctcaagtcac agttaaatag cccgatactt tcctcgctca 540  
 ctatgagcta ttacagcata catttttagta ctacatactt attcagtaaa aagccctcaa 600  
 aattgaagac aaaggacggg atccccgggt accgagctcg aattcaggcc tctagatctc 660  
 attattcctc catcaagaga agctccacgc tgtccacgat gaaggttccc tcggtttcac 720  
 cgatctcgat ccacactttg tcggtctcag gaaagtactc aagctccttg gtaacatagc 780  
 caactggaag tgggtgtgtag tccctgtaac ctctgttgaa ctcgcaaggg ttctcacgtc 840  
 tgccatctgt gtaggatttc tcctcgtaac cggaggcata gtcagcagga acggaaggag 900  
 cttcggtgta acctctgtta cggctagtgt aggcacctcc gtactcttcc tgattcacag 960  
 tgtagtcggt gcaagtaacg gtgttggttg gatagatttc ttctcgacg cagttggaga 1020  
 acttaagctc gtcggtgttg ttctcgatct cgtggatggc cacgcaacct tcaccgtatc 1080  
 cctccttgta agcggtcaca cggagaatgt agcctctacc tggacagact ctaacctctt 1140  
 gggacacttc agcttccac tcaggcaca ccaggacgga acgctgattg ttctgttctt 1200  
 ccacgtccac atgaccttcc acattccagc agctgaggcc attggtgaag tcaccgttct 1260  
 tgatgacgtt tctggcatcg tacaaggaga atgcggtaaa gatacgtccc tcaagttcct 1320  
 cgaagatggc agcgttcaca ccagggatca cggacaactc aggcaagtaa gcttcacgaa 1380  
 tgctgtgcac acgtttgtct gcggcggtga tcatggcgat gttggtgtcg gcttgcaact 1440  
 gatcatattg ggagttcacg aacaaagcat ccacggactc tttggcctcc ttgtaaacga 1500

## 52288B-SEQ-listing.txt

tgtagtttc ccattcgagt ttctcacgtt tgtccctcca cttcttctct gctctcttca	1560
cacgagcgag agcttcaccg accaatgggt tctcttcgag aaactcaagg ttgccaagtc	1620
ttgcgtgtcc gtcttgggtc ttgatcttga agatgacca gactccgagg tcctcattca	1680
ggtcagtaca tcccacatcg atgtccaagg agaagtgatg agaatgggtg gcacacttct	1740
cgccatccct gcaggagcag tccaagtcag gattccactc aagggtgtgga gcgcattctgt	1800
taggctctcc acacttccca atgggagatt gggcagaaag tggccagagg gaaccagtac	1860
ctgggacatt cacggtctcg tgcttggcat tgtacctgat cgagtagatt tcaaggctct	1920
ggctgtcttc gatgtagcct ctaagttgat acctggtgaa ggctttgagt ttggactcat	1980
cgatcttctg gtacaagtag gtagggtagc actcgtcgaa agttccggag agggtgacgt	2040
agttctcctt gaacacatcg tcgctcctt ggatgggtgat cccggtgctt ccacccaac	2100
cacgttctgg ctgcctgttg atgtcttga agttggagtc ttgcaagaga ttcctctcgt	2160
cgctgagacg cttggcgtgt ttaactttct cggagagttc acgcttctcg tcgaggcaga	2220
actcatcgct aaggtaggtg accaagttgg aacttgggtc aatgtgatag tcagtaacgt	2280
tagttttcaa gccaagctga ttgggtggagg taaagagggc gttcacagcc ttctgggctc	2340
tctcaagggt gtactcagcc tcgagtgttg cagtaactgg aatgaactcg aatctgtcga	2400
taatcactcc tgcagtccca ctaaagtttc taacaccac gatgttaccg agtgaagatg	2460
taaaagcatt ggcactttca aagtaaccga aatcgctgga ttggagatta tccaaggagg	2520
tagctgtagc tggaactgta ttggagaaga tggatgaatt accccaatta acgttgaggt	2580
gaataggggt cacagaagca tacctcacac gaactctata tctggtagat gtggatggga	2640
agtgaattgg aacttcaata taccctctat tctgaatgtt atttccactg ctgttgagtc	2700
taacgaggtc tccaccagtg aatcctgggtc ctgaaatgac agaaccgttg aagagaaagt	2760
ttcccttcac tgcagggatt tgagtaatac tatcggtatg gatgatgttg ttgaactcag	2820
cactacgatg tatccaagag aacataggag ctctgatgat gctcacggaa ctgttgctga	2880
atccggaacg gaacatggac acgtgggtca acctgtggga gaatccttgc ctgggtggca	2940
cattgttggt ctgtggaaaa ggaaggtggc tcctacaaat gccatcattg cgataaagga	3000
aaggccatcg ttgaagatgc ctctgccgac agtgggtcca aagatggacc cccaccacg	3060
aggagcatcg tggaaaaaga agacgttcca accacgtctt caaagcaagt ggattgatgt	3120
gatatctcca ctgacgtaag ggatgacgca caatcccact atccttcgca agacccttc	3180
tctatataag gaagttcatt tcatttggag aggacacgct gacaagctga ctctagcaga	3240
tctccatgga caacaacca aacatcaacg aatgcattcc atacaactgc ttgagtaacc	3300
cagaagtga agtacttggg ggagaacgca ttgaaaccgg ttacactccc atcgacatct	3360
ccttgctcctt gacacagttt ctgctcagcg agttcgtgcc aggtgctggg ttcgttctcg	3420

## 52288B-SEQ-listing.txt

```

gactagttga catcatctgg ggtatctttg gtccatctca atgggatgca ttcttggtgc 3480
aaattgagca gttgatcaac cagaggatcg aagagttcgc caggaaccag gccatctcta 3540
ggttggaagg attgagcaat ctctaccaa tctatgcaga gagcttcaga gagtggaag 3600
ccgatcctac taaccagct ctccgcgagg aaatgcgtat tcaattcaac gacatgaaca 3660
gcgccctgac cacagctatc ccattgttcg cagtccagaa ctaccaagtt cctctcttgt 3720
ccgtgtacgt tcaagcagct aatcttcacc tcagcgtgct tcgagacggt agcgtgtttg 3780
ggcaaagggt gggattcgat gctgcaacca tcaatagccg ttacaacgac cttactaggc 3840
tgattgga aa ctacaccgac cacgctgttc gttggtacaa cactggcttg gagcgtgtct 3900
ggggtcctga ttctagagat tggattagat acaaccagtt caggagagaa ttgacctca 3960
cagttttgga cattgtgtct ctcttccga actatgactc cagaacctac cctatccgta 4020
cagtgtccca acttaccaga gaaatctata ctaaccagtt tcttgagaac ttcgacggt 4080
gcttccgtgg ttctgccaa ggtatcgaag gctccatcag gagccacac ttgatggaca 4140
tcttgaacag cataactatc tacaccgatg ctacagagg agagtattac tggcttgac 4200
accagatcat ggcctctcca gttggattca gcgggccga gtttaccttt cctctctatg 4260
gaactatggg aaacgccgct ccacaacaac gtatcgttg tcaactaggt cagggtgtct 4320
acagaacaaa cactgatagt ttaaactgaa ggcgggaaac gacaatctga tcccagcttg 4380
catgcctgca ggtcaattca atattgtggc aggacattgc tacatgatac ctcttagaat 4440
tgtttagact tcagatcgat cttgtcagtc tgaaagaccc aaaaacaaat gcaatttctt 4500
ttctggtaga ccgtgacaat ttgtctaaga tgtatctgat ttaatgcctt ttgtatataa 4560
tacactcadc taatctagtt aatttagctt cagagtaa at tacttcagca tatttatacg 4620
tgccaagtgc caaccatadc aaattagcta agcagacagt tgaagtacac aaaacaaaag 4680
catcatatgc tgatttattt attcatagat ggagctcaag tcatagttaa atagcccgat 4740
actttcctcg ctactatga gctattacag catacatttt agtactacat acttattcag 4800
taaaaagccc taaaattga agacaaagga cgggatcccc ggggtaccgag ctggaattca 4860
ggcctctaga tctcattatt cctccatcaa gagaagctcc acgctgtcca cgatgaaggt 4920
tccctcggtt tcaccgatct cgatccacac tttgtcggtc tcaggaaagt act 4973

```

```

<210> 16
<211> 19
<212> DNA
<213> artificial sequence

```

```

<220>
<221> misc_feature
<222> (1)..(19)
<223> 5' primer to 5' flanking sequence of SEQ ID NO: 15 from 8 to 26;

```



```

<400> 16
gacttcccat cttctatcc
19

<210> 17
<211> 19
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(19)
<223> 3'primer to partial e35S promoter of SEQ ID NO: 15 from 3154 to 3
136

<400> 17
attgtgcgtc atcccttac
19

<210> 18
<211> 22
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(22)
<223> 5'primer to partial 3' cry1Ac sequence of SEQ ID NO: 15 from 2581
to 2603

<400> 18
gaataggggt cacagaagca ta
22

<210> 19
<211> 20
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(20)
<223> 3'primer to partial 5' cry1Ac sequence of SEQ ID NO: 15 from 3455
to 3435

<400> 19
ggaccaaaga taccccagat
20

<210> 20
<211> 19
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(19)
<223> 5'primer to partial e35S promoter of SEQ ID NO: 15 from 2993 to 3
011

```

<400> 20  
ataaaggaaa ggccatcgt 19

<210> 21  
<211> 25  
<212> DNA  
\* <213> artificial sequence  
  
<220>  
^<221> misc\_feature  
<222> (1)..(25)  
<223> 3'primer to full-length cry1Ac sequence of SEQ ID NO: 15 from 497  
3 to 4949

<400> 21  
agtactttcc tgagaccgac aaagt 25

10